



SEQUENCE LISTING

B/
<110> Donoho, Gregory
Turner, C. Alexander Jr.
Nehls, Michael
Friedrich, Glenn
Zambrowicz, Brian
Sands, Arthur T.

<120> Novel Human Proteins and Polynucleotides
Encoding the Same

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<150> US 60/160,285

<151> 1999-10-19

<150> US 60/183,583

<151> 2000-02-18

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TECH CENTER 1600/2900

MAY 06 2002

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Phe Thr Ser Ser Ser Asp Gln Tyr Gly Pro Tyr Cys Gly Ser Met Thr
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Val Pro Lys Glu Leu Leu Leu Asn Thr Ser Glu Val Thr Val Arg Phe
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Ser Ser Asp His Pro Asp Leu Ile Thr Cys Leu Glu Arg Ala Ser His
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Tyr Leu Lys Thr Glu Tyr Ser Lys Phe Cys Pro Ala Gly Cys Arg Asp
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165 170 175
Glu Gly Ile Leu Ala Asn Gly Val Leu Ser Arg Asp Gly Ser Leu Ser
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Gln Asp Gln Gly Pro Ser Trp Ala Ser Gly Asp Ser Ser Asn Asn His
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Lys Pro Arg Glu Trp Leu Glu Ile Asp Leu Gly Glu Lys Lys Ile
260 265 270
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Leu Val Trp Arg Lys Thr Ser Gln Ser Thr Ser Val Ser Thr Lys Lys
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Glu Asp Glu Thr Ile Thr Arg Pro Ile Pro Ser Glu Glu Thr Ser Thr
385 390 395 400

Gly Ile Asn Ile Thr Thr Val Ala Ile Pro Leu Val Leu Val Val
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gcccgggc	tgggtatgg	ctgtggacac	ctagtgactt	atcaggatag	tggcacaatg	300
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gatttaataa	catgtttg	acgagct	cattatttga	agacagaata	cagcaaattc	660
tgc	gtttagaga	cgtac	gacatttctg	ggaatatgtt	agatggat	720
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ggtgc	gatc	tttga	gac	ttc	ccat	960
cagtc	tttga	tttga	tttgc	tttgc	tttgc	1020
gaccaagg	tttgc	tttgc	tttgc	tttgc	tttgc	1080
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aacttcc	tttgc	tttgc	tttgc	tttgc	tttgc	1320
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actaaga	tttgc	tttgc	tttgc	tttgc	tttgc	1500
ataaaacat	tttgc	tttgc	tttgc	tttgc	tttgc	1560
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Ala Glu Glu Leu Gly Asp Gly Cys Gly His Leu Val Thr Tyr Gln Asp
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Tyr Leu Leu Phe Thr Ser Ser Asp Gln Tyr Gly Pro Tyr Cys Gly
145 150 155 160
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Val Arg Phe Glu Ser Gly Ser His Ile Ser Gly Arg Gly Phe Leu Leu
180 185 190
Thr Tyr Ala Ser Ser Asp His Pro Asp Leu Ile Thr Cys Leu Glu Arg
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Ala Ser His Tyr Leu Lys Thr Glu Tyr Ser Lys Phe Cys Pro Ala Gly
210 215 220
Cys Arg Asp Val Ala Gly Asp Ile Ser Gly Asn Met Val Asp Gly Tyr
225 230 235 240
Arg Asp Thr Ser Leu Leu Cys Lys Ala Ala Ile His Ala Gly Ile Ile
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Ala Asp Glu Leu Gly Gln Ile Ser Val Leu Gln Arg Lys Gly Ile
260 265 270
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370 375 380
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385 390 395 400
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Gln Gly Asn Ser Asn Phe Arg Asp Pro Val Gln Asn Asn Phe Ile Pro
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Pro Ile Val Ala Arg Tyr Val Arg Val Val Pro Gln Thr Trp His Gln
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 465 470 475 480
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 485 490 495
 Thr Ser Thr Gly Ile Asn Ile Thr Thr Val Ala Ile Pro Leu Val Leu
 500 505 510
 Leu Val Val Leu Val Phe Ala Gly Met Gly Ile Phe Ala Ala Phe Arg
 515 520 525
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 530 535 540
 Thr Asp Cys Trp Lys Gln Ile Lys Tyr Pro Phe Ala Arg His Gln Ser
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tggaaagcaga	ttaaatatcc	ctttgccaga	aggctcagaa	tttggatggat	1560
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Gly Ser Met Thr Val Pro Lys Glu Leu Leu Asn Thr Ser Glu Val
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145 150 155 160
Arg Ala Ser His Tyr Leu Lys Thr Glu Tyr Ser Lys Phe Cys Pro Ala
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Gly Cys Arg Asp Val Ala Gly Asp Ile Ser Gly Asn Met Val Asp Gly
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Tyr Arg Asp Thr Ser Leu Leu Cys Lys Ala Ala Ile His Ala Gly Ile
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275 280 285
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 485 490 495
 Lys Thr Asp Cys Trp Lys Gln Ile Lys Tyr Pro Phe Ala Arg His Gln
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